

# SEQUENCE LISTING

<110> LINDNER, Volkhard  
FRIESEL, Robert F.

<120> COMPOSITIONS, METHODS AND KITS RELATING TO REMODEL

<130> 36-13 Lindner et al. (200036.0013)

<140> Not Yet Assigned

<141> 2000-10-19

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<170> PatentIn Ver. 2.1

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| Pro         | Gly | Arg | Asp | Gly | Phe | Lys | Gly | Glu | Lys | Gly | Glu | Cys | Leu | Arg | Glu |
|             |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     |     | 95  |
| Ser         | Phe | Glu | Glu | Ser | Trp | Thr | Pro | Asn | Tyr | Lys | Gln | Cys | Ser | Trp | Ser |
|             |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Ser         | Leu | Asn | Tyr | Gly | Ile | Asp | Leu | Gly | Lys | Ile | Ala | Glu | Cys | Thr | Phe |
|             |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Thr         | Lys | Met | Arg | Ser | Asn | Ser | Ala | Leu | Arg | Val | Leu | Phe | Ser | Gly | Ser |
|             |     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |
| Leu         | Arg | Leu | Lys | Cys | Arg | Asn | Ala | Cys | Cys | Gln | Arg | Trp | Tyr | Phe | Thr |
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|             |     |     | 165 |     |     |     |     | 170 |     |     |     |     |     | 175 |     |
| Tyr         | Leu | Asp | Gln | Gly | Ser | Pro | Glu | Met | Asn | Ser | Thr | Ile | Asn | Ile | His |
|             |     | 180 |     |     |     |     |     | 185 |     |     |     |     | 190 |     |     |
| Arg         | Thr | Ser | Ser | Val | Glu | Gly | Leu | Cys | Glu | Gly | Ile | Gly | Ala | Gly | Leu |
|             |     | 195 |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |
| Val         | Asp | Val | Ala | Ile | Trp | Val | Gly | Thr | Cys | Ser | Asp | Tyr | Pro | Lys | Gly |
|             | 210 |     |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |
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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
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| Met | His | Pro | Gln | Gly | Arg | Ala | Ala | Ser | Pro | Gln | Leu | Leu | Leu | Gly | Leu |  |
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| Phe | Leu | Val | Leu | Leu | Leu | Leu | Leu | Gln | Leu | Ser | Ala | Pro | Ser | Ser | Ala |  |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |  |
| Ser | Glu | Asn | Pro | Lys | Val | Lys | Gln | Lys | Ala | Leu | Ile | Arg | Gln | Arg | Glu |  |
| 65  |     |     |     |     | 70  |     |     |     |     | 75  |     |     |     |     | 80  |  |
| Val | Val | Asp | Leu | Tyr | Asn | Gly | Met | Cys | Leu | Gln | Gly | Pro | Ala | Gly | Val |  |
|     |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     | 95  |     |  |
| Pro | Gly | Arg | Asp | Gly | Ser | Pro | Gly | Ala | Asn | Gly | Ile | Pro | Gly | Thr | Pro |  |
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| Trp | Ser | Ser | Leu | Asn | Tyr | Gly | Ile | Asp | Leu | Gly | Lys | Ile | Ala | Glu | Cys |  |
| 145 |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     |     | 160 |  |
| Thr | Phe | Thr | Lys | Met | Arg | Ser | Asn | Ser | Ala | Leu | Arg | Val | Leu | Phe | Ser |  |
|     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |     |  |
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| Phe | Thr | Phe | Asn | Gly | Ala | Glu | Cys | Ser | Gly | Pro | Leu | Pro | Ile | Glu | Ala |  |
|     | 195 |     |     |     |     |     | 200 |     |     |     |     | 205 |     |     |     |  |
| Ile | Ile | Tyr | Leu | Asp | Gln | Gly | Ser | Pro | Glu | Leu | Asn | Ser | Thr | Ile | Asn |  |
|     | 210 |     |     |     |     | 215 |     |     |     |     | 220 |     |     |     |     |  |
| Ile | His | Arg | Thr | Ser | Ser | Val | Glu | Gly | Leu | Cys | Glu | Gly | Ile | Gly | Ala |  |
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| Gly | Leu | Val | Asp | Val | Ala | Ile | Trp | Val | Gly | Thr | Cys | Ser | Asp | Tyr | Pro |  |
|     |     | 245 |     |     |     |     |     | 250 |     |     |     |     | 255 |     |     |  |
| Lys | Gly | Asp | Ala | Ser | Thr | Gly | Trp | Asn | Ser | Val | Ser | Arg | Ile | Ile | Ile |  |
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15

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20

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&lt;211&gt; 734

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

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&lt;223&gt; Description of Artificial Sequence:myc-tagged

REMODEL construct

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734